AMENDMENTS TO THE CLAIMS

This listing of claims will replace all prior versions, and listings, of claims in the application:

Listing of Claims:

- 1. (Original) A process for the enhanced production of pantothenate, comprising culturing a microorganism having a deregulated methylenetetrahydrofolate (MTF) biosynthetic pathway, under conditions such that pantothenate production is enhanced.
- 2. (Original) A process for the enhanced production of pantothenate, comprising culturing a microorganism having
 - (i) a deregulated pantothenate biosynthetic pathway, and
- (ii) a deregulated methylenetetrahydrofolate (MTF) biosynthetic pathway, under conditions such that pantothenate production is enhanced.
- 3. (Original) The process of claim 2, wherein said microorganism has at least two pantothenate biosynthetic enzymes deregulated.
- 4. (Original) The process of claim 2, wherein said microorganism has at least three pantothenate biosynthetic enzymes deregulated.
- 5. (Original) The process of claim 2, wherein said microorganism has at least four pantothenate biosynthetic enzymes deregulated.
- 6. (Original) The process of claim 5, wherein said microorganism has a deregulated ketopantoate hydroxymethyltransferase, a deregulated ketopantoate reductase, a deregulated pantothenate synthetase and a deregulated aspartate-α-decarboxylase.
- 7. (Currently Amended) The process of any one of claims 1 and 3-6 elaims 1 to 6, wherein said microorganism further has a deregulated isoleucine-valine (*ilv*) biosynthetic pathway.
- 8. (Original) The process of claim 7, wherein said microorganism has at least two isoleucine-valine (*ilv*) biosynthetic enzymes deregulated.

9. (Original) The process of claim 7, wherein said microorganism has at least three isoleucine-valine (*ilv*) biosynthetic enzymes deregulated.

- 10. (Original) The process of claim 9, wherein said microorganism has a deregulated acetohydroxyacid acid synthetase, a deregulated acetohydroxyacid isomeroreductase, and a deregulated dihydroxyacid dehydratase.
- 11. (Original) The process of any one of claims 1 to 10, wherein the microorganism has at least one MTF biosynthetic enzyme deregulated.
- 12. (Original) The process of claim 11, wherein the microorganism has a deregulated *glyA* gene.
- 13. (Original) The process of claim 11, wherein the microorganism has a deregulated *serA* gene.
- 14. (Original) The process of claim 11, wherein the microorganism has a deregulated glyA gene and a deregulated serA gene.
- 15. (Currently Amended) The process of claim 12 or 14, wherein the <u>glyA</u> gene is deregulated by mutating, deleting, or disrupting a the microorganism has a mutated, deleted or disrupted purR gene in said microorganism.
- 16. (Currently Amended) The A-process of claim 2, wherein for the production pantothenate, comprising culturing a microorganism further has having a deregulated pantothenate biosynthetic pathway, a deregulated isoleucine-valine (*ilv*) biosynthetic pathway, and a deregulated methylenetetrahydrofolate (MTF) biosynthetic pathway deregulated,
- 17. (Currently Amended) The A-process of claim 16, wherein for the production pantothenate, comprising culturing a microorganism having a deregulated pantothenate biosynthetic pathway, a deregulated isoleucine valine (ilv) biosynthetic pathway, and a deregulated methylenetetrahydrofolate (MTF) biosynthetic pathway, is cultured under conditions such that at least 50 g/L pantothenate is produced after 36 hours of culturing the microorganism.

18. (Currently Amended) The process of claim 17, comprising culturing the microorganism <u>under conditions</u> such that at least 60 g/L pantothenate is produced after 36 hours of culturing the microorganism.

- 19. (Currently Amended) The process of claim 17, comprising culturing the microorganism under conditions such that at least 70 g/L pantothenate is produced after 36 hours of culturing the microorganism.
- 20. (Currently Amended) The A-process of claim 16, for the production pantothenate, comprising culturing the a microorganism having a deregulated pantothenate biosynthetic pathway, a deregulated isoleucine valine (ilv) biosynthetic pathway, and a deregulated methylenetetrahydrofolate (MTF) biosynthetic pathway deregulated, under conditions such that at least 60 g/L pantothenate is produced after 48 hours of culturing the microorganism.
- 21. (Currently Amended) The process of claim 20, comprising culturing the microorganism under conditions such that at least 70 g/L pantothenate is produced after 48 hours of culturing the microorganism.
- 22. (Currently Amended) The process of claim 20, comprising culturing the microorganism <u>under conditions</u> such that at least 80 g/L pantothenate is produced after 48 hours of culturing the microorganism.
- 23. (Currently Amended) The process of any one of the preceding claims 1, 2, and 16, wherein pantothenate production is further enhanced by regulating pantothenate kinase activity.
- 24. (Original) The process of claim 23, wherein pantothenate kinase activity is decreased.
- 25. (Original) The process of claim 24, wherein CoaA is deleted and CoaX is downregulated.
- 26. (Original) The process of claim 24, wherein CoaX is deleted and CoaA is downregulated.

27. (Original) The process of claim 24, wherein CoaX and CoaA are downregulated.

- 28. (Currently Amended) The process of any one of the above claims 1, 2, and 16, wherein said microorganism is cultured under conditions of excess serine.
 - 29. (Canceled)
- 30. (Currently Amended) The process of any one of the above claims 1, 2, wherein said microorganism has the pantothenate biosynthetic pathway deregulated such that pantothenate production is independent of β -alanine feed.
- 31. (Currently Amended) The process of any one of the above claims 1, 2, wherein the microorganism is a Gram positive microorganism.
- 32. (Currently Amended) The process of any one of the above claims 1, 2, wherein the microorganism belongs to the genus *Bacillus*.
- 33. (Currently Amended) The process of any one of the above claims 1, 2, wherein the microorganism is *Bacillus subtilis*.
 - 34.-41. (Canceled)
- 42. (Currently Amended) The A-process of claim 16, for producing pantothenate comprising culturing a recombinant wherein said microorganism has having:
 - (a) a deregulated panB gene; and
 - (b)—a deregulated panD gene; and
 - (c) at least one deregulated isoleucine-valine (ilv) biosynthetic enzymeencoding gene;

and is cultured under conditions such that at least 30 g/l pantothenate is produced after 36 hours of culturing the microorganism.

43. (Currently Amended) The process of claim 42, wherein said microorganism further has a deregulated methylenetetrahydrofolate (MTF) biosynthetic pathway

and said microorganism is cultured under conditions such that at least 50 g/l pantothenate is produced after 36 hours of culturing the microorganism.

44.-45. (Canceled)

- 46. (Currently Amended) <u>The A process of claim 2, for producing</u> pantothenate comprising culturing a recombinant wherein said microorganism has having:
 - (a)—a deregulated panB gene;
 - (b)—a deregulated panD gene; and
 - (c)—a deregulated glyA gene;

<u>and is cultured</u> under conditions of excess valine, such that at least 50 g/l pantothenate is produced after 36 hours of culturing the microorganism.

- 47. (Currently Amended) <u>The A process of claim 2, for producing</u>

 pantothenate comprising culturing a recombinant wherein said microorganism has having:
 - (a)—a deregulated panB gene;
 - (b)—a deregulated panD gene; and
 - (e)—a mutated, deleted or disrupted purR gene;

and is cultured under conditions of excess valine, such that at least 50 g/l pantothenate is produced after 36 hours of culturing the microorganism.

- 48. (Currently Amended) <u>The A process of claim 2, for producing pantothenate comprising culturing a recombinant wherein said microorganism has having:</u>
 - (a)—a deregulated panB gene;
 - (b)—a deregulated panD gene; and
 - (c)—a deregulated serA gene;

and is cultured under conditions of excess valine, such that at least 50 g/l pantothenate is produced after 36 hours of culturing the microorganism.

- 49. (Currently Amended) <u>The A process of claim 2, for producing</u> pantothenate comprising culturing a recombinant wherein said microorganism has having:
 - (a)—a deregulated panB gene;
 - (b)—a deregulated panD gene;
 - (e) a deregulated serA gene; and
 - (d) —a deregulated glyA gene; and

is cultured under conditions of excess valine, such that at least 50 g/l pantothenate is produced after 36 hours of culturing the microorganism.

- 50. (New) The process of any one of claims, 1, 2, 7, 16, 23, 28, 30, and 42, wherein the microorganism has a deregulated glyA gene.
- 51. (New) The process of claim 50, wherein the glyA gene is deregulated by mutating, deleting, or disrupting a purR gene in said microorganism.